STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/597,/93
Source:	1FWO,
Date Processed by STIC:	8/24/06
•	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
 U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
 Alexandria, VA 22314

Revised 01/10/06



IFWO

RAW SEQUENCE LISTING DATE: 08/24/2006 TIME: 15:19:46 PATENT APPLICATION: US/10/597,173

Input Set : F:\APP SER NO 10597173 (sequence list).txt

Output Set: N:\CRF4\08242006\J597173.raw

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1 <110> APPLICANT: GENEIN CO., LTD.
            KIM, Cheol-Min
             PARK, Hee-Kyung
     5 <120> TITLE OF INVENTION: Oligonucleotide for genotyping of Mycoplasma, microarray
             comprising the oligonucleotide, and method for detection of
             species using the microarray
     9 <130> FILE REFERENCE: PN053079
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/597,173
C--> 11 <141> CURRENT FILING DATE: 2006-07-13
                                                                    Does Not Comply
     11 <160> NUMBER OF SEQ ID NOS: 133
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    15 <210> SEO ID NO: 1
    16 <211> LENGTH: 344
    17 <212> TYPE: DNA
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    27 atatttacaa cgacatcaaa aatcaaatta atggttaatt tgttttgatt catcgagtaa
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RAW SEQUENCE LISTING DATE: 08/24/2006
PATENT APPLICATION: US/10/597,173 TIME: 15:19:46

Input Set : F:\APP SER NO 10597173 (sequence list).txt
Output Set: N:\CRF4\08242006\J597173.raw

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	74 atttcaaagt ttagatcaac ctatagaata caaaatcaat acaataggtc aatactatac	180
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	89 gcctaaaagt cttatatcta gttttgagag gacatcctct ctaattgttc tttgaaaact	180
	91 gaatagtaaa ttttttgata tttacaacga catctaaata attgaattaa gtcaatttgt	240
	93 ttagatttca tcgagatagt cattttaaaa aaatgattca ttgaaatgtc ttaaaataca	300
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	101 <213> ORGANISM: M.synoviae 103 <400> SEQUENCE: 6	
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	108 agagaacaac ctctcttaaa attgttcttt gaaaactaaa tagtaataaa gatattacaa	180
	110 cgacatcaaa aatataaatt aattaaggtt aatttgtttt gataccgagt ttaaattatt	240
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Input Set : F:\APP SER NO 10597173 (sequence list).txt
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 RAW SEQUENCE LISTING
 DATE: 08/24/2006

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Input Set : F:\APP SER NO 10597173 (sequence list).txt

Output Set: N:\CRF4\08242006\J597173.raw

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Input Set : F:\APP SER NO 10597173 (sequence list).txt

Output Set: N:\CRF4\08242006\J597173.raw

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VARIABLE LOCATION SUMMARY

DATE: 08/24/2006 TIME: 15:19:47

PATENT APPLICATION: US/10/597,173

Input Set : F:\APP SER NO 10597173 (sequence list).txt

Output Set: N:\CRF4\08242006\J597173.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:8; N Pos. 20

VERIFICATION SUMMARYDATE: 08/24/2006PATENT APPLICATION: US/10/597,173TIME: 15:19:47

Input Set : F:\APP SER NO 10597173 (sequence list).txt

Output Set: N:\CRF4\08242006\J597173.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:140 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:8 L:140 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:8

L:140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0